

Seq. search result attached

LOCUS ATF18O22 100269 bp DNA linear PLN 13-APR-2000
DEFINITION Arabidopsis thaliana DNA chromosome 5, BAC clone F18O22 (ESSA project).
ACCESSION AL163817
VERSION AL163817.1 GI:7573446
KEYWORDS
SOURCE Arabidopsis thaliana (thale cress)
ORGANISM Arabidopsis thaliana
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
REFERENCE 1 (bases 1 to 100269)
AUTHORS Bevan,M., Murphy,G., Ridley,P., Hudson,S., Bancroft,I., Mewes,H.W.,
Rudd,S., Lemcke,K. and Mayer,K.F.X.
JOURNAL Unpublished
REFERENCE 2 (bases 1 to 100269)
AUTHORS EU Arabidopsis sequencing,project.
TITLE Direct Submission
JOURNAL Submitted (13-APR-2000) MIPS, at the Max-Planck-Institut fuer
Biochemie, Am Klopferspitz 18a, D-82152 Martinsried, FRG, E-mail:
lemcke@mips.biochem.mpg.de,mayer@mips.biochem.mpg.de Project
Coordinator: Mike Bevan, Molecular Genetics Department, Cambridge
Laboratory, John Innes Centre, Colney Lane, NR4 7UJ Norwich, UK,
E-mail: michael.bevan@bbsrc.ac.uk
COMMENT Information on performance of analysis and a more detailed
annotation of this entry and other sequences of chromosomes 3, 4
and 5 can be viewed at: <http://www.mips.biochem.mpg.de/proj/thal/>.
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CAB87779. putative protein ...[gi:7573465]

LOCUS CAB87779 382 aa linear PLN 13-APR-2000

DEFINITION putative protein [Arabidopsis thaliana].

ACCESSION CAB87779

VERSION CAB87779.1 GI:7573465

DBSOURCE embl locus ATF18O22, accession AL163817.1

KEYWORDS

SOURCE Arabidopsis thaliana (thale cress)

ORGANISM Arabidopsis thaliana

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

REFERENCE 1 (residues 1 to 382)

AUTHORS Bevan,M., Murphy,G., Ridley,P., Hudson,S., Bancroft,I., Mewes,H.W.,
Rudd,S., Lemcke,K. and Mayer,K.F.X.

JOURNAL Unpublished

REFERENCE 2 (residues 1 to 382)

AUTHORS EU Arabidopsis sequencing,project.

TITLE Direct Submission

JOURNAL Submitted (13-APR-2000) MIPS, at the Max-Planck-Institut fuer
Biochemie, Am Klopferspitz 18a, D-82152 Martinsried, FRG, E-mail:
lemcke@mips.biochem.mpg.de,mayer@mips.biochem.mpg.de Project
Coordinator: Mike Bevan, Molecular Genetics Department, Cambridge
Laboratory, John Innes Centre, Colney Lane, NR4 7UJ Norwich, UK,
E-mail: michael.bevan@bbsrc.ac.uk

COMMENT Information on performance of analysis and a more detailed
annotation of this entry and other sequences of chromosomes 3, 4
and 5 can be viewed at: <http://www.mips.biochem.mpg.de/proj/thal/>.

FEATURES Location/Qualifiers

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